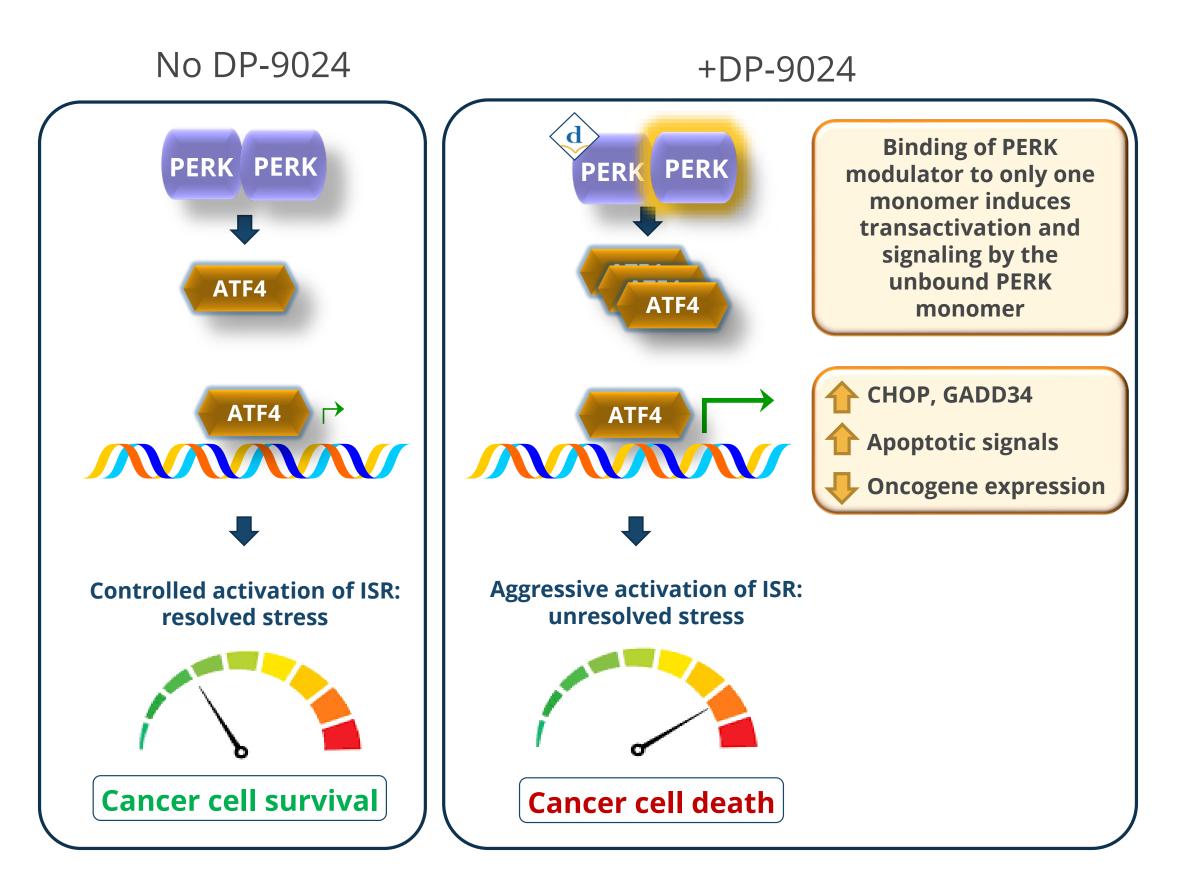
DP-9024, an investigational small molecule modulator of the integrated stress response kinase PERK, causes B-cell cancer growth inhibition as single agent and in combination with standard-of-care agents

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Introduction

- The Integrated Stress Response (ISR) is a major adaptive stress response pathway in cancer cell maintenance¹⁻⁴
- The ISR family member PERK is a member of the Unfolded Protein Response (UPR) pathway which plays a role in resolving endoplasmic reticulum (ER) stress such as processing unfolded proteins¹⁻⁴
- The UPR is considered an Achilles' heel in B-cell cancers, as multiple myeloma and B-cell lymphoma are dependent on a well-balanced UPR pathway to cope with the high demand for protein folding and their secretory nature
- Given the double-edged-sword nature of the UPR, the activation of PERK and its downstream pathway can have cytoprotective or cytotoxic effects⁵
- In B-cell cancers, the UPR is at close-to-maximum cytoprotective capacity, such that further pharmacological stimulation of PERK can potentially be leveraged to cause a cancer cell cytotoxic response and induce antitumoral effects



Methods

- Modulation of ISR kinases was characterized using enzymatic assays
- Kinome selectivity profiling was determined using enzymatic and cellular assays
- Cellular modulation of the ISR/UPR pathway (PERK, ATF4, and CHOP) or the apoptosis pathway (c-PARP, c-Caspase 3/7) was measured by Western blot, qRT-PCR, or ELISA
- *In vivo* upregulation of tumoral ATF4 was determined in a MM pharmacokinetic/pharmacodynamic xenograft model
- *In vivo* inhibition of tumor growth was determined in MM and B-cell lymphoma xenografts

Results

DP-9024 was designed as a selective and potent modulator of ISR kinases that activates PERK, with optimized pharmaceutical and selectivity profiles

	DP-9024	
Cellular assays	PERK dimerization assay (BRET; EC ₅₀ , nM)	274
	H929 ATF4 stimulation (fold increase versus control)	12
Off-target profile	Kinome and safety	Highly selective
	hERG (Predictor [™] fluorescence polarization; IC ₂₀ , μM)	>20
ADME	Microsomal stability (human, mouse) % remaining at 60 min	64%, 70%
	Caco-2 (A-B, efflux ratio)	41, 1.6

DP-9024 exhibits a selective kinome profile

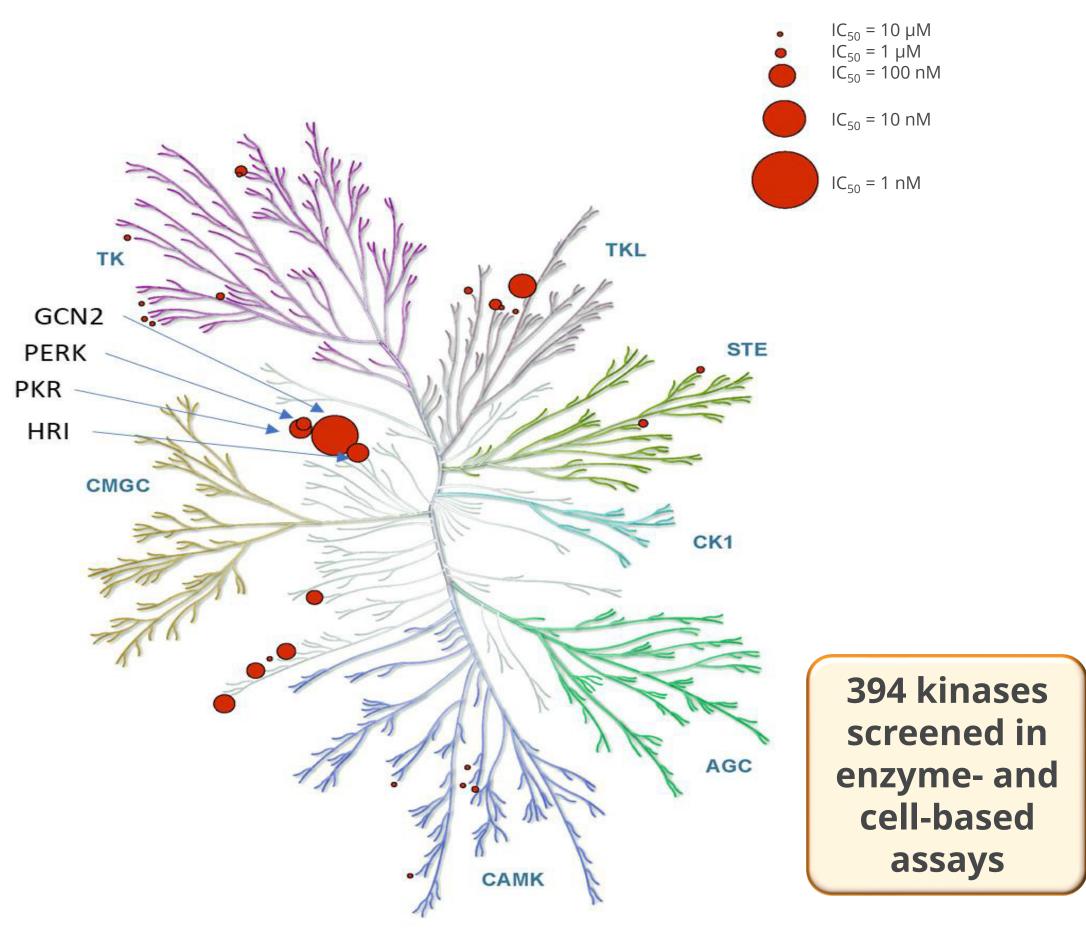


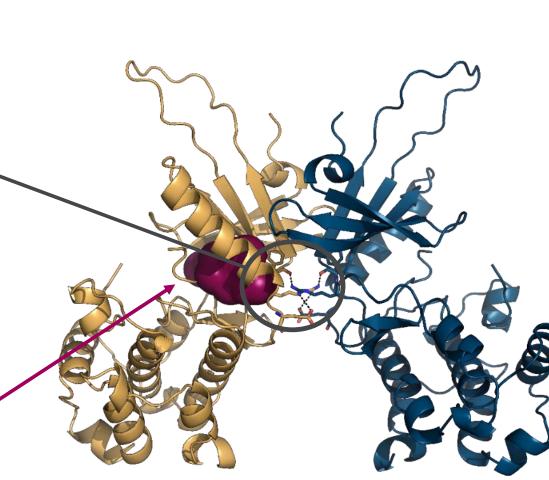
Illustration reproduced courtesy of Cell Signaling Technology, Inc.

Structure of close DP-9024 analog bound to PERK monomer

PERK forms a head-to-head dimer nucleated by transmonomer hydrogen bond interactions with key arginine residues in the **C-helix** switch

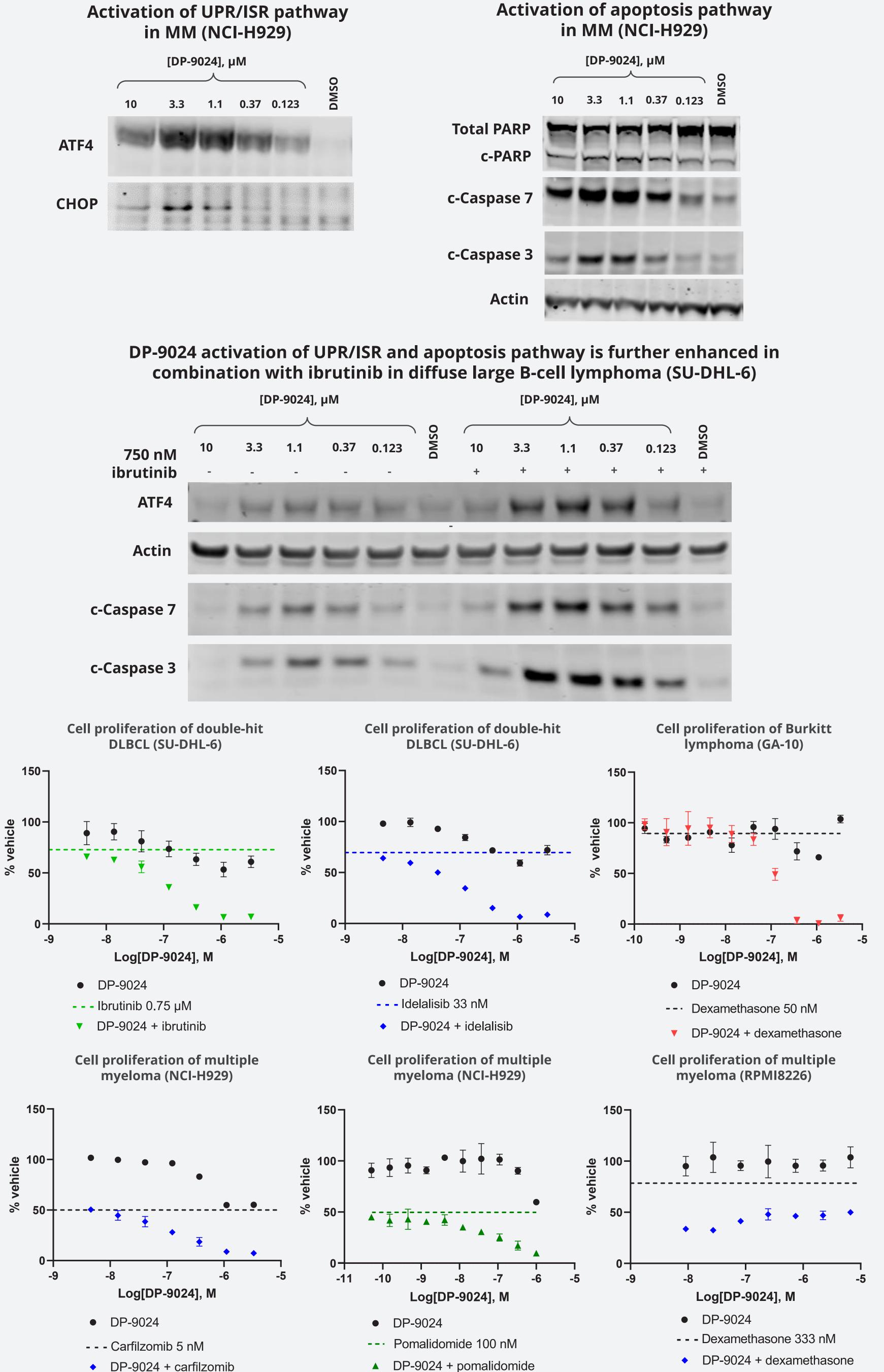
Binding **C-helix switch** in the "C-helix out" state limits inhibitor binding to half of the dimer, **transactivating** the unbound PERK half of the dimer in the catalytically active, **"C-helix in"** state

PRESENTED AT THE AMERICAN ASSOCIATION FOR CANCER RESEARCH (AACR) ANNUAL MEETING ORLANDO, FL, APRIL 14–19, 2023



Both monomers were bound to compound under the crystal structure-saturating conditions. In this figure, only 1 bound monomer is shown to represent model of activation at or below IC_{50} of DP-9024 binding.

PERK activator DP-9024 activates the ISR pathway, induces apoptosis, and inhibits B-cell proliferation in combination with clinical agents



ACKNOWLEDGMENTS

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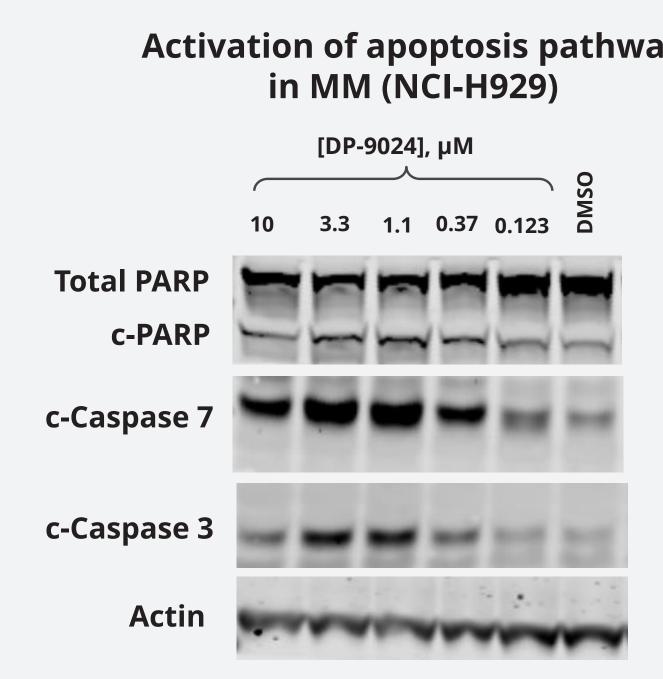
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Gada Al-Ani (Galani@Deciphera.com) All authors are/were full time employees of Deciphera Pharmaceuticals, LLC and own/owned Deciphera Pharmaceuticals, LLC

CORRESPONDING

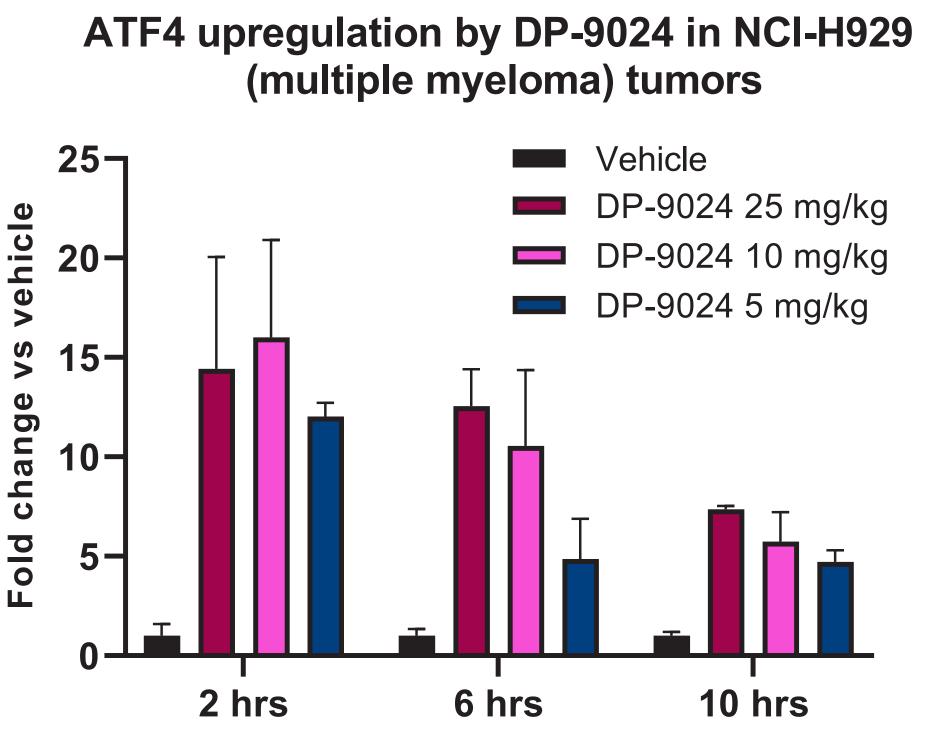
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AUTHOR/DISCLOSURES



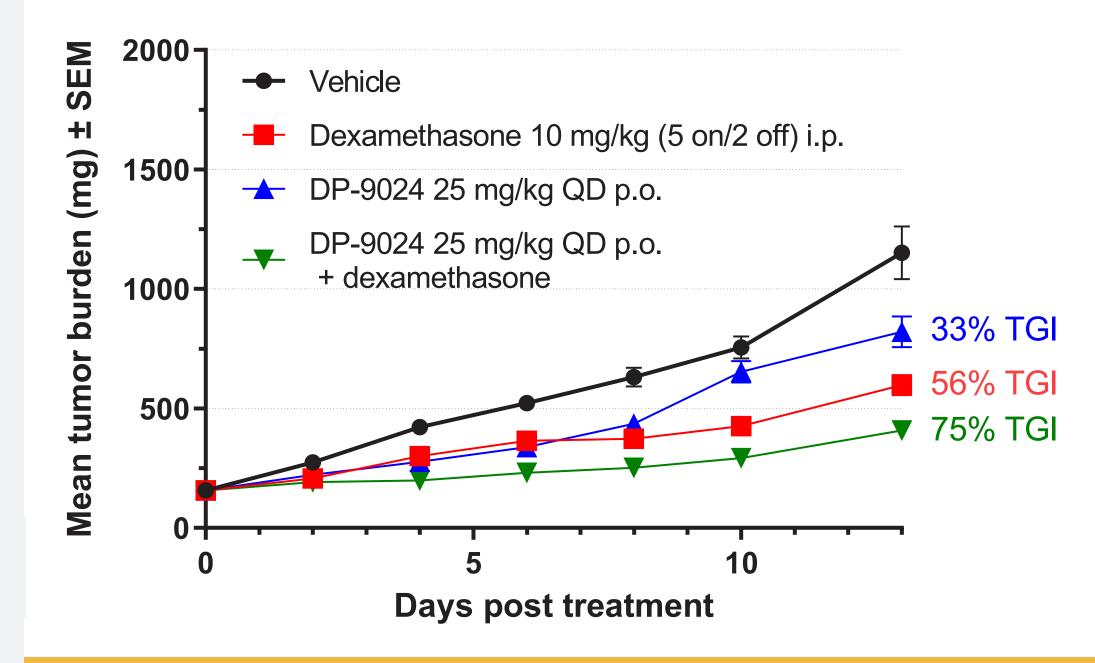
DP-9024 + dexamethasone

DP-9024 upregulates tumoral ISR and inhibits B-cell cancer tumor growth in combination with anti-myeloma agents in mouse xenograft models in vivo



Dose (mg/kg)	Timepoint (hrs)	DP-9024 free plasma levels (ng/mL)	Fold ATF4 upregulation
25	2	99	14
25	6	104	13
25	10	116	7
10	2	70	16
10	6	47	11
10	10	59	6
5	2	42	12
5	6	38	5
5	10	48	5

CA46 (Burkitt lymphoma) tumor growth



CONCLUSIONS

ABBREVIATIONS

ADME, absorption, distribution, metabolism, and excretion; ADP, adenosine diphosphate; AGC, protein kinase A, G, and C families; ATF4, activating transcription factor 4 BRET, bioluminescence resonance energy transfer; c-, cleaved; CHOP, C/EBP homologous protein; CK1, casein kinase 1 family; CMGC, family of kinases including cyclin-

dependent kinases, mitogen-activated protein kinases, glycogen synthase kinases, and cyclin-dependent kinases; DLBCL, diffuse large B-cell lymphoma; DMSO, dimethyl

reaction; s.c., subcutaneous; SEM, standard error of the mean; STE, homologs of yeast sterile 7, sterile 11, and sterile 20 kinase family; TGI, tumor growth inhibition; TK,

tyrosine kinase family; TKL, tyrosine kinase-like family; UPR, unfolded protein response; VEGFA, vascular endothelial growth factor A.

sulfoxide; EC₅₀, half maximal effective concentration; ELISA, enzyme-linked immunosorbent assay; ER, endoplasmic reticulum; GADD34, growth arrest and DNA damage-

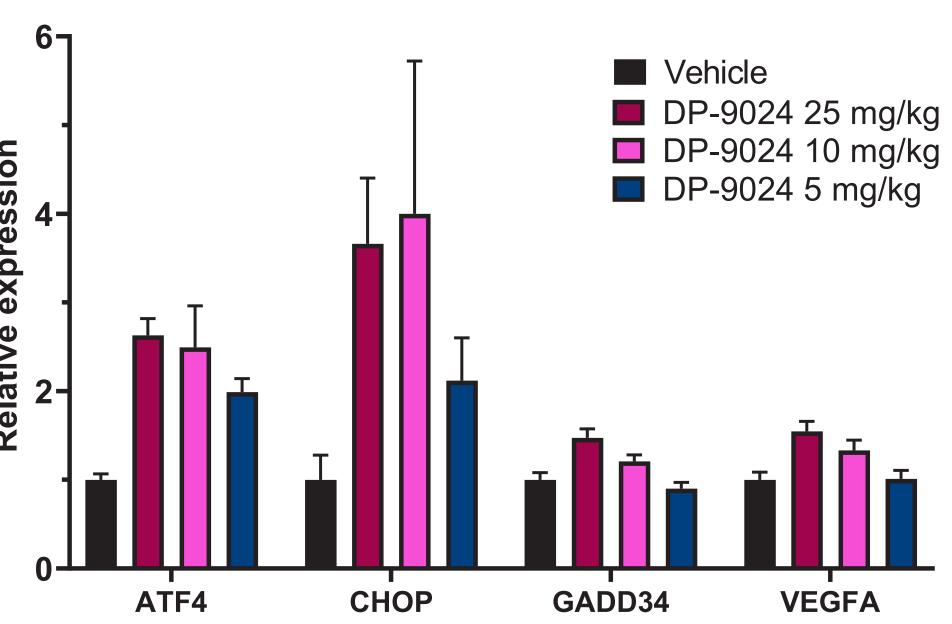
inducible protein 34; GCN2, general control nonderepressible 2; hERG, human ether-a-go-go-related gene; HRI, heme-regulated inhibitor; IC₂₀, concentration inducing 20%

inhibition; IC₅₀, half maximal inhibitory concentration; i.p., intraperitoneally; ISR, integrated stress response; MM, multiple myeloma; PARP, poly ADP-ribose polymerase; PER

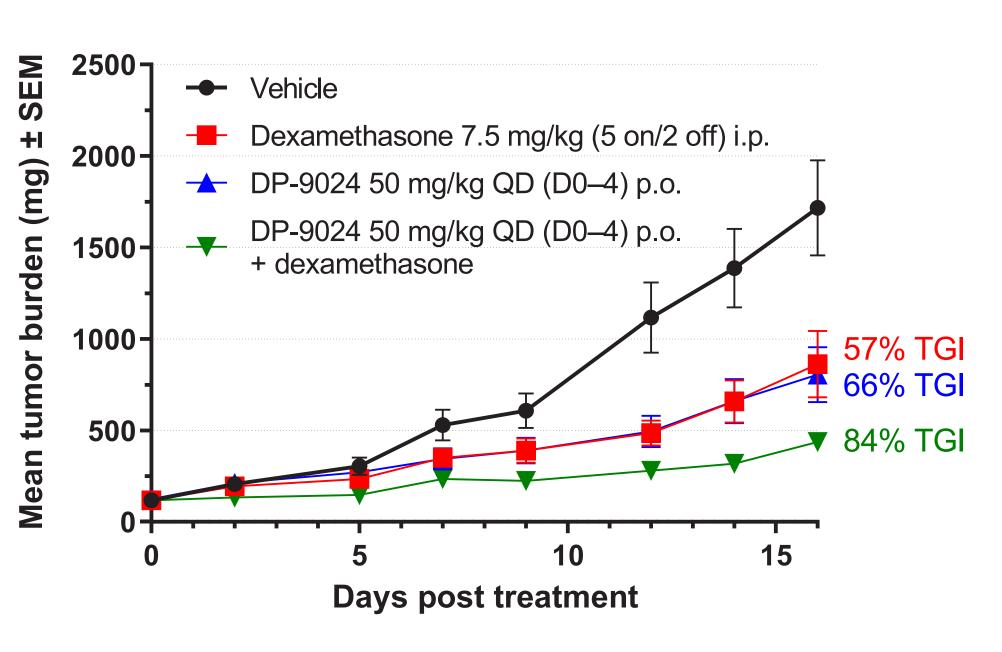
protein kinase R-like endoplasmic reticulum kinase; PKR, protein kinase R; p.o., orally; QD, once daily; qRT-PCR, real-time quantitative reverse-transcription polymerase chain

deciphera® **Poster: 1640**

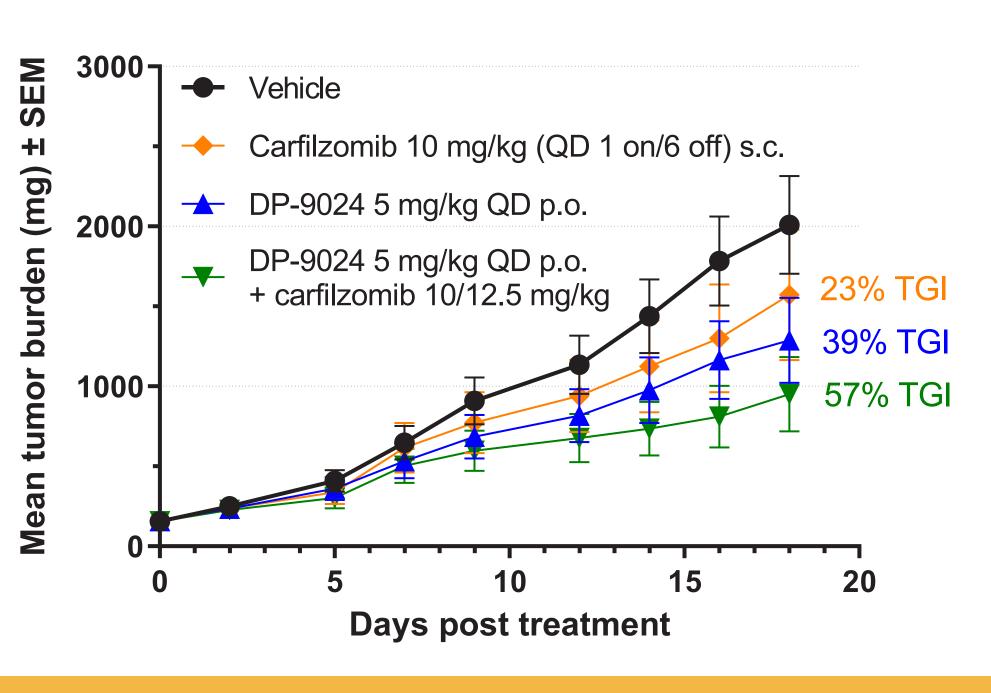
Upregulation of ISR genes by DP-9024 in NCI-H929 (multiple myeloma) tumors



RPMI8226 (multiple myeloma) tumor growth



RPMI8226 (multiple myeloma) tumor growth



• The ISR/UPR is a targetable vulnerability in cancers with high basal levels of ER stress • DP-9024 increases UPR signaling by directly binding to one PERK monomer and inducing dimerization and transactivation of the unbound PERK monomer

• This novel mechanism leads to antitumoral effects in B-cell cancers *in vitro* and *in vivo*, likely through the induction of unresolved UPR stress, which may provide an alternative mechanism to current UPR-targeting therapies

> REFERENCES 1. Marciniak SJ, et al. Nat Rev Drug Dlscov. 2022;21(2):115-40 2. Pakos-Zebrucka K, et al. *EMBO Rep*. 2016;17(10):1374-95. 3. Licari E, et al. Int | Biochem Cell Biol. 2021;139:106059. 4. White-Gilbertson S, et al. *Front Genet*. 2013;4:109.

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